SEQUENCE LISTING

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<110> Ghayur, Tarig et al.
<120> ANTIBODIES THAT BIND HUMAN INTERLEUKIN-18 AND METHODS
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Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile 35 40 45

Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile 50 55 60

Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile 65 70 75 80

Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
85 90 95

Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
100 105 110

Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu 115 120 125

Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu 130 135 140

Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp 150 155

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Gly Gln Asp Met Glu Gln Gln Val Val Phe Ser Met Ser Phe Val Gln 35 40 45

Gly Glu Glu Ser Asn Asp Lys Ile Pro Val Ala Leu Gly Leu Lys Glu 50 55 60

Lys Asn Leu Tyr Leu Ser Cys Val Leu Lys Asp Asp Lys Pro Thr Leu 65 70 75 80

Gln Leu Glu Ser Val Asp Pro Lys Asn Tyr Pro Lys Lys Met Glu 85 90 95

Lys Arg Phe Val Phe Asn Lys Ile Glu Ile Asn Asn Lys Leu Glu Phe 100 105 110

Glu Ser Ala Gln Phe Pro Asn Trp Tyr Ile Ser Thr Ser Gln Ala Glu 115 120 125

Asn Met Pro Val Phe Leu Gly Gly Thr Lys Gly Gly Gln Asp Ile Thr 130 135 140

Asp Phe Thr Met Gln Phe Val Ser Ser 145 150

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Asn Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro His
Ala Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val
Lys Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr
Asp Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile Arg
Ser Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly
                                105
Trp Phe Leu Cys Thr Ala Met Glu Ala Asp Gln Pro Val Ser Leu Thr
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Asp
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Leu Lys His Cys Ser Cys Ser Leu Ala His Glu Ile Glu Thr Thr
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Lys Ser Trp Tyr Lys Ser Ser Gly Ser Gln Glu His Val Glu Leu Asn
Pro Arg Ser Ser Arg Ile Ala Leu His Asp Cys Val Leu Glu Phe
Trp Pro Val Glu Leu Asn Asp Thr Gly Ser Tyr Phe Phe Gln Met Lys
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Asn Tyr Thr Gln Lys Trp Lys Leu Asn Val Ile Arg Arg Asn Lys His

85 90 Ser Cys Phe Thr Glu Arg Gln Val Thr Ser Lys Ile Val Glu Val Lys 105 Lys Phe Phe Gln Ile Thr Cys Glu Asn Ser Tyr Tyr Gln Thr Leu Val Asn Ser Thr Ser Leu Tyr Lys Asn Cys Lys Leu Leu Leu Glu Asn 135 Asn Lys Asn Pro Thr Ile Lys Lys Asn Ala Glu Phe Glu Asp Gln Gly Tyr Tyr Ser Cys Val His Phe Leu His His Asn Gly Lys Leu Phe Asn 170 Ile Thr Lys Thr Phe Asn Ile Thr Ile Val Glu Asp Arg Ser Asn Ile 185 Val Pro Val Leu Gly Pro Lys Leu Asn His Val Ala Val Glu Leu Gly Lys Asn Val Arg Leu Asn Cys Ser Ala Leu Leu Asn Glu Glu Asp 215 Val Ile Tyr Trp Met Phe Gly Glu Glu Asn Gly Ser Asp Pro Asn Ile 230 235 His Glu Glu Lys Glu Met Arg Ile Met Thr Pro Glu Gly Lys Trp His Ala Ser Lys Val Leu Arg Ile Glu Asn Ile Gly Glu Ser Asn Leu Asn Val Leu Tyr Asn Cys Thr Val Ala Ser Thr Gly Gly Thr Asp Thr Lys Ser Phe Ile Leu Val Arg Lys Ala Asp <210> 8 <211> 310 <212> PRT <213> Homo sapiens <400> 8

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Lys Glu Gly Ala
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Gly Lys Asn Asn Arg Pro Ser
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Gly Ser Arg Asp Ser Ser Gly Ile His Val Val
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Gln Gly Asp Ser Leu Arg His Phe Tyr Ser Asn
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Gln
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Gln
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                                     10
Ser Met Lys Val Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Gly Tyr
                                 25
Tyr Ile His Trp Val Arg Gln Ala His Gly Gln Gly Phe Glu Trp Ile
                             40
Gly Arg Leu Asn Pro Thr Thr Gly Asp Ala Asn Phe Ala Glu Lys Phe
                         55
Gln Gly Arg Val Ala Leu Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
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Leu Gln Leu Asp Ser Leu Lys Ser Asp Asp Thr Ala Val Tyr Tyr Cys
Ala Gly Lys Glu Gly Ala Trp Gly Gln Gly Thr Leu Val Thr Val Ser
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Asn Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
Gly Lys Asn Asn Arg Pro Ser Gly Ile Pro Asp Arg Phe Ser Gly Ser
Gly Ser Gly Asn Thr Gly Ser Leu Thr Ile Thr Gly Ala Gln Ala Glu
Asp Glu Ala Asp Tyr Tyr Cys Gly Ser Arg Asp Ser Ser Gly Ile His
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Val Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu Gly
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Asp Asp Asp Tyr Asp Phe Asp Tyr
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Ala Ala Trp Asp Asp Ser Leu Ser Gly Pro Val
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Gly
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Ala Ile Ser Gly Ser Gly Gly Ser Thr Trp Tyr Ala Asp Ser Val Lys
Gly
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<213> Homo sapiens
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 Phe Thr Phe Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly
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20

 Lys Gly Leu Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Ser Thr 35 40 45

Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn 50 55 60

Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp 65 70 75 80

Thr Ala Val Tyr Tyr Cys Ala Arg Asp Asp Asp Asp Tyr Asp Phe Asp 85 90 95

Tyr Trp Gly Arg Gly Thr Met Val Thr Val Ser Ser 100 105

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Arg Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Ile Asn 20 25 30

Ala Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu 35 40 45

Ile Tyr Gly Asn Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser 50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
65 70 75 80

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu 85 90 95

Ser Gly Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
100 105 110

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Phe Thr Phe Ser Ser Tyr Ala Met Ser Trp Val Arg Gln Ala Pro Gly 20 25 30

Lys Gly Leu Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr 35 40 45

Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Asp Asp Tyr Asp Phe Asp Tyr Trp Gly Arg Gly Thr Met Val Thr Val Ser Ser Gly Gly Gly 105 Ser Gly Gly Gly Ser Gly Gly Gly Ser Ala Gln Ser Val Leu Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln Arg Val Thr Ile 135 Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Ile Asn Ala Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr Gly Asn 170 Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Lys Ser 185 Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln Ser Glu Asp Glu 195 200 Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu Ser Gly Pro Val 215 Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly 230 <210> 31 <211> 14 <212> PRT <213> Homo sapiens Tyr Phe Gly Lys Leu Glu Ser Lys Leu Ser Val Ile Arg Asn <210> 32 <211> 14 <212> PRT <213> Homo sapiens <400> 32

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       <211> 14
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       <213> Homo sapiens
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       <400> 36
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113
       Asn Arg Pro Leu Phe Glu Asp Met Thr Asp Ser Asp Cys Arg
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       <212> PRT
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       <211> 14
       <212> PRT
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       <213> Homo sapiens
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       Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
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       <212> PRT
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       Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp
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   <212> PRT
   <213> Homo sapiens
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Asp Gln Val Leu Phe Ile Asp Gln Gly Asn Arg Pro Leu Phe Glu Asp
Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
                                105
Met Gln Phe Glu Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
        115
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tcg atg aaa gtc tcc tgt aag act tct gga tac acc ttc acc ggc tat
                                                                   96
Ser Met Lys Val Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Gly Tyr
                                 25
tat atc cac tgg gtg cga cag gcc cct gga cag gga ttc gag tgg ata
                                                                   144
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Phe Glu Trp Ile
gga cgg ctc aac ccc acc act ggt gac gca aat ttt gca gaa aag ttt
                                                                   192
Gly Arg Leu Asn Pro Thr Thr Gly Asp Ala Asn Phe Ala Glu Lys Phe
cag age agg ate ace eta ace aga ace ace tee ate age aca gee tat
                                                                   240
Gln Gly Arg Val Ala Leu Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
tta caa cta gac agc ctc aaa tct gac gac acg gcc gta tat tat tgt
                                                                   288
Leu Gln Leu Asp Ser Leu Lys Ser Asp Asp Thr Ala Val Tyr Tyr Cys
geg gga aaa gag ggt gee tgg gge cag gge ace etg gte ace gte teg
                                                                   336
Ala Gly Lys Glu Gly Ala Trp Gly Gln Gly Thr Leu Val Thr Val Ser
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                                105
agt gg
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Ser
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Ser Met Lys Val Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Gly Tyr
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Thr Val Arg Ile Thr Cys Gln Gly Asp Ser Leu Arg His Phe Tyr Pro 20 25 30

Asn Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr 35 40 45

Gly Lys Asn Asn Arg Pro Ser Gly Ile Pro Asp Arg Phe Ser Gly Ser 50 55 60

Gly Ser Gly Asn Thr Gly Ser Leu Thr Ile Thr Gly Ala Gln Ala Glu 65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gly Ser Arg Asp Ser Ser Gly Ile His 85 90 95

Val Val Phe Gly Gly Gly Thr Lys Val Thr Val Leu Gly 100 105

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1 5 10 15

tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttt agc agc tat 96 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr 20 25 30

gcc atg agc tgg gtc cgc cag gct cca ggg aag ggg ctg gag tgg gtc 144
Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

tca gct att agt ggt agt ggt agc aca tac tac gca gac tcc gtg 192 Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val 50 55

aag ggc cgg ttc acc atc tcc aga gac aat tcc aag aac acg ctg tat 240 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr 65 70 75 80

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gcg aga gat gac gat gac tac gac ttt gac tac tgg ggc cgg ggg aca 336 Ala Arg Asp Asp Asp Tyr Asp Phe Asp Tyr Trp Gly Arg Gly Thr

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